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## AMENDMENTS TO THE CLAIMS

The following listing of claims replaces all prior versions, and listings, of claims in this application.

## Listing of Claims

1. (Currently amended) A composition comprising a protein in crystalline form wherein the protein consists of SEQ. ID No. 4 SEQ ID NO.4, and wherein the protein crystal has a crystal lattice in a  $P2_12_12_1$  space group and unit cell dimensions,  $\pm/-5\%$ , of a=48.36Å b=72.29Å and c=94.52Å,  $\alpha=\beta=\gamma=90^\circ$ .

2-3 (Cancelled)

4. (Previously presented) A composition according to claim 1 wherein the protein crystal diffracts X-rays for a determination of structure coordinates to a resolution having a value that is less than or equal to 3.0 Angstroms.

5-8 (Cancelled)

9. (Currently amended) A method for forming a crystal of a protein comprising:

forming a crystallization volume comprising a precipitant solution and a protein that consists of SEQ. ID No. 4 SEQ ID NO.4, and wherein a protein crystal is formed that has a crystal lattice in a  $P2_{12}1_{21}$  space group and unit cell dimensions, +/- 5%, of a=48.36Å b=72.29Å and c=94.52Å,  $\alpha$ = $\beta$ = $\gamma$ =90°; and

storing the crystallization volume under conditions suitable for formation of a protein crystal.

10-11. (Cancelled)

12. (Previously presented) A method according to claim 9 wherein a protein crystal is formed that diffracts X-rays for a determination of structure coordinates to a resolution having a value that is less than or equal to 3.0 Angstroms.

13-14. (Canceled)

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15. (Previously presented) A method according to claim 9, wherein a protein crystal is formed, the method further comprising diffracting the protein crystal to produce a diffraction pattern and solving the structure of the protein from the diffraction pattern.

16-17, (Cancelled)

18. (Withdrawn – currently amended) A method of identifying an entity that associates with a protein comprising:

taking structure coordinates from diffraction data obtained from a crystal of a protein that has at least 90% identity with SEQ. ID No. 3 consists of SEQ ID NO.3, and wherein the protein crystal has a crystal lattice in a P2<sub>1</sub>2<sub>1</sub>2<sub>1</sub> space group and having unit cell dimensions,  $\pm$ 7.5%, of a=48.36Å b=72.29Å and c=94.52Å,  $\pm$ 8=7.29°; and

performing rational drug design using a three dimensional structure that is based on the obtained structure coordinates.

19-21. (Canceled)

- 22. (Withdrawn) A method according to claim 18, the method further comprising selecting one or more entities based on the rational drug design and contacting the selected entities with the protein.
- 23. (Withdrawn) A method according to claim 18, the method further comprising measuring an activity of the protein when contacted with the one or more entities.
- 24. (Withdrawn) A method according to claim 18, the method further comprising comparing activity of the protein in a presence of and in the absence of the one or more entities; and selecting entities where activity of the protein changes depending whether a particular entity is present.
- 25. (Withdrawn) A method according to claim 18, the method further comprising contacting cells expressing the protein with the one or more entities and detecting a change in a phenotype of the cells when a particular entity is present.
- 26. (Withdrawn) The method according to claim 15 further comprising:

performing rational drug design using the solved structure; and

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identifying an entity that associates with the protein.

- 27. (Withdrawn) The method according to claim 26 further comprising selecting one or more entities based on the rational drug design and contacting the selected entities with the protein.
- 28. (Withdrawn) The method according to claim 27 further comprising measuring an activity of the protein when contacted with the one or more entities.
- (Currently amended) A composition comprising a soluble protein consisting of SEQ. ID No.4
  SEQ ID NO.4.
- 30. (Withdrawn currently amended) A e<del>omposition comprising a soluble protein consisting of SEQ. ID No. 3 SEQ ID NO. 3.</del>
- 31. (New) An isolated soluble protein consisting of residues 136-461 of SEQ ID NO:1.
- (New) An isolated non-crystalline protein consisting of residues 136-461 of SEQ ID NO:1.
- 33. (New) A non-crystalline protein consisting of SEQ ID NO:4.
- 34. (New-Withdrawn) A non-crystalline protein consisting of SEQ ID NO:3.
- 35. (New) A composition comprising a protein in crystalline form wherein the protein consists of residues 136-461 of SEQ ID NO:1, and wherein the protein crystal has a crystal lattice in a  $P2_12_12_1$  space group and unit cell dimensions, +/- 5%, of a=48.36Å b=72.29Å and c=94.52Å,  $\alpha=\beta=\gamma=90^\circ$ .
- 36. (New) A method for forming a crystal of a protein comprising:

forming a crystallization volume comprising a precipitant solution and a protein that consists of residues 136-461 of SEQ ID NO:1, and wherein a protein crystal is formed that has a crystal lattice in a  $P2_12_12_1$  space group and unit cell dimensions, +/- 5%, of a=48.36Å b=72.29Å and c=94.52Å,  $\alpha$ = $\beta$ = $\gamma$ =90°; and

storing the crystallization volume under conditions suitable for formation of a protein crystal.